

=> file medline hcaplus biosis biotechds scisearch embase  
COST IN U.S. DOLLARS SINCE FILE TOTAL  
ENTRY SESSION  
FULL ESTIMATED COST 0.21 0.21

FILE 'MEDLINE' ENTERED AT 16:56:45 ON 29 JUL 2004

FILE 'HCAPLUS' ENTERED AT 16:56:45 ON 29 JUL 2004  
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=> s dimethylallyl-cycloacetoacetyl-L-tryptophan synthase  
L-1 1 DIMETHYLLALLYL-CYCLOACETOACETYL-L-TRYPTOPHAN SYNTHASE

=> d 11 ibib ab

toxin-deficient *Aspergillus* sp. (61pp)

=> file registry		SINCE FILE	TOTAL
COST IN U.S. DOLLARS		ENTRY	SESSION
FULL ESTIMATED COST		10.46	10.67

FILE 'REGISTRY' ENTERED AT 16:57:37 ON 29 JUL 2004  
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Property values tagged with IC are from the ZIC/VINITI data file  
provided by InfoChem.

STRUCTURE FILE UPDATES: 28 JUL 2004 HIGHEST RN 718597-29-6  
DICTIONARY FILE UPDATES: 28 JUL 2004 HIGHEST RN 718597-29-6

TSCA INFORMATION NOW CURRENT THROUGH MAY 21, 2004

Please note that search-term pricing does apply when  
conducting SmartSELECT searches.

Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. For more  
information enter HELP PROP at an arrow prompt in the file or refer  
to the file summary sheet on the web at:  
<http://www.cas.org/ONLINE/DBSS/registryss.html>

=> s dimethylallyl-cycloacetoacetyl-L-tryptophan synthase  
810 DIMETHYLALLYL  
2 CYCLOACETOACETYL  
1568416 L  
34465 TRYPTOPHAN  
25801 SYNTHASE  
L2 0 DIMETHYLALLYL-CYCLOACETOACETYL-L-TRYPTOPHAN SYNTHASE  
(DIMETHYLALLYL (W) CYCLOACETOACETYL (W) L (W) TRYPTOPHAN (W) SYNTHASE)

=> s dimethylallyl-cycloacetoacetyl tryptophan synthase  
810 DIMETHYLALLYL  
2 CYCLOACETOACETYL  
34465 TRYPTOPHAN  
25801 SYNTHASE  
L3 1 DIMETHYLALLYL-CYCLOACETOACETYL TRYPTOPHAN SYNTHASE  
(DIMETHYLALLYL (W) CYCLOACETOACETYL (W) TRYPTOPHAN (W) SYNTHASE)

=> d 13

L3 ANSWER 1 OF 1 REGISTRY COPYRIGHT 2004 ACS on STN  
RN 280784-57-8 REGISTRY  
CN DNA (*Aspergillus oryzae* strain A1560 gene DCAT-S  
dimethylallylcycloacetoacetyltryptophan synthase cDNA plus flanks)  
(9CI) (CA INDEX NAME)

OTHER NAMES:

CN 1: PN: WO0039322 SEQID: 1 claimed DNA  
FS NUCLEIC ACID SEQUENCE  
MF Unspecified  
CI MAN  
SR CA  
LC STN Files: CA, CAPLUS, TOXCENTER, USPATFULL  
DT.CA CAplus document type: Patent  
RL.P Roles from patents: BIOL (Biological study); OCCU (Occurrence); PROC  
(Process); PRP (Properties)

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*  
\*\*\* USE 'SQD' OR 'SQIDE' FORMATS TO DISPLAY SEQUENCE \*\*\*  
1 REFERENCES IN FILE CA (1907 TO DATE)  
1 REFERENCES IN FILE CAPLUS (1907 TO DATE)

=> s dimethylallylcycloacetoacetyltryptophan synthase  
2 DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN  
25801 SYNTHASE  
L4 1 DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE  
(DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN (W) SYNTHASE)

COST IN U.S. DOLLARS	SINCE FILE ENTRY	TOTAL SESSION
FULL ESTIMATED COST	52.18	62.85

FILE 'MEDLINE' ENTERED AT 16:59:46 ON 29 JUL 2004

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=> s dimethylallylcycloacetoacetyltryptophan synthase  
L5 1 DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE

=> d 15 ibib ab

L5 ANSWER 1 OF 1 HCAPLUS COPYRIGHT 2004 ACS on STN  
ACCESSION NUMBER: 2000:457232 HCAPLUS  
DOCUMENT NUMBER: 133:85094  
TITLE: Protein production in *Aspergillus* mutant cells that  
produce decreased levels of toxin  
INVENTOR(S): Christensen, Bjorn Eggert; Mollgaard, Henrik;  
Kaasgaard, Svend; Lehmbeck, Jan  
PATENT ASSIGNEE(S): Novo Nordisk A/S, Den.  
SOURCE: PCT Int. Appl., 66 pp.  
CODEN: PIXXD2  
DOCUMENT TYPE: Patent  
LANGUAGE: English  
FAMILY ACC. NUM. COUNT: 1  
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2000039322	A1	20000706	WO 1999-DK726	19991222
W:	AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA,			

MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI,  
 SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ,  
 BY, KG, KZ, MD, RU, TJ, TM  
 RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE,  
 DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF,  
 CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG  
 EP 1141371 A1 20011010 EP 1999-960956 19991222  
 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,  
 IE, SI, LT, LV, FI, RO  
 JP 2002533133 T2 20021008 JP 2000-591212 19991222  
 US 6383781 B1 20020507 US 1999-472364 19991223  
 US 2002197682 A1 20021226 US 2002-99704 20020315  
 PRIORITY APPLN. INFO.: DK 1998-1726 A 19981223  
 DK 1999-745 A 19990527  
 US 1999-117396P P 19990127  
 US 1999-139593P P 19990617  
 WO 1999-DK726 W 19991222  
 US 1999-472364 A3 19991223

AB A method is provided for producing a polypeptide of interest by (a) cultivating a mutant of a parent *Aspergillus* cell, wherein (i) the mutant comprises a first nucleic acid sequence encoding the polypeptide and a second nucleic acid sequence comprising a modification of at least one of the genes responsible for the biosynthesis or secretion of at least one toxin, and (ii) the mutant produces less of the toxin than the parent *Aspergillus* cell when cultured under the same conditions; and (b) isolating the polypeptide from the culture medium. Also, mutants of *Aspergillus* cells are provided, as well as methods for obtaining the mutant cells. Thus, the dimethylallylcycloacetoacetyl-L-tryptophan synthase gene (DCAT-S) was characterized in *Aspergillus oryzae*. Since this enzyme is involved in biosynthesis of cyclopiazonic acid (CPA), CPA-neg. strains were constructed by modification of the DCAT-S gene, and improved of lipase and xylanase genes was obsd. in the CPA-neg. strains. Strains lacking genes *aflR* and *omtA*, involved in aflatoxin biosynthesis, also demonstrated improved heterologous protein prodn.

REFERENCE COUNT: 4 THERE ARE 4 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

=> dimethylallylcycloacetoacetyl synthase  
 DIMETHYLALLYLCYCLOACETOACETYL IS NOT A RECOGNIZED COMMAND  
 The previous command name entered was not recognized by the system.  
 For a list of commands available to you in the current file, enter  
 "HELP COMMANDS" at an arrow prompt (=>).

=> s dimethylallylcycloacetoacetyl synthase  
 L6 0 DIMETHYLALLYLCYCLOACETOACETYL SYNTHASE

=> s dimethylallyl cycloacetoacetyl l-tryptophan synthase  
 L7 0 DIMETHYLALLYL CYCLOACETOACETYL L-TRYPTOPHAN SYNTHASE

=> s *Aspergillus oryzae* and synthase  
 L8 125 ASPERGILLUS ORYZAE AND SYNTHASE

=> dup rem 18  
 PROCESSING COMPLETED FOR L8  
 L9 62 DUP REM L8 (63 DUPLICATES REMOVED)

=> s 19 and dimethylallylcycloacetoacetyltryptophan synthase  
 L10 1 L9 AND DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE

=> d his

(FILE 'HOME' ENTERED AT 16:56:06 ON 29 JUL 2004)

FILE 'MEDLINE, HCAPLUS, BIOSIS, BIOTECHDS, SCISEARCH, EMBASE' ENTERED AT

16:56:45 ON 29 JUL 2004

L1 1 S DIMETHYLALLYL-CYCLOACETOACETYL-L-TRYPTOPHAN SYNTHASE

FILE 'REGISTRY' ENTERED AT 16:57:37 ON 29 JUL 2004

L2 0 S DIMETHYLALLYL-CYCLOACETOACETYL-L-TRYPTOPHAN SYNTHASE

L3 1 S DIMETHYLALLYL-CYCLOACETOACETYL TRYPTOPHAN SYNTHASE

L4 1 S DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE

FILE 'MEDLINE, HCAPLUS, BIOSIS, BIOTECHDS, SCISEARCH, EMBASE' ENTERED AT  
16:59:46 ON 29 JUL 2004

L5 1 S DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE

L6 0 S DIMETHYLALLYLCYCLOACETOACETYL SYNTHASE

L7 0 S DIMETHYLALLYL CYCLOACETOACETYL L-TRYPTOPHAN SYNTHASE

L8 125 S ASPERGILLUS ORYZAE AND SYNTHASE

L9 62 DUP REM L8 (63 DUPLICATES REMOVED)

L10 1 S L9 AND DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE

=> s l9 and tryptophan synthase  
L11 2 L9 AND TRYPTOPHAN SYNTHASE

=> d l11 1-2 ibib ab

L11 ANSWER 1 OF 2 MEDLINE on STN

ACCESSION NUMBER: 96358132 MEDLINE

DOCUMENT NUMBER: PubMed ID: 8762144

TITLE: Invariant glycines and prolines flanking in loops the  
strand beta 2 of various (alpha/beta)8-barrel enzymes: a  
hidden homology?.

AUTHOR: Janecek S

CORPORATE SOURCE: Institute of Ecobiology, Slovak Academy of Sciences,  
Bratislava, Slovakia.. sjanecek@ue.savba.sk

SOURCE: Protein science : a publication of the Protein Society,  
(1996 Jun) 5 (6) 1136-43.  
Journal code: 9211750. ISSN: 0961-8368.

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 199612

ENTRY DATE: Entered STN: 19970128

Last Updated on STN: 19980206

Entered Medline: 19961213

AB The question of parallel (alpha/beta)8-barrel fold evolution remains  
unclear, owing mainly to the lack of sequence homology throughout the  
amino acid sequences of (alpha/beta)8-barrel enzymes. The "classical"  
approaches used in the search for homologies among (alpha/beta)8-barrels  
(e.g., production of structurally based alignments) have yielded  
alignments perfect from the structural point of view, but the approaches  
have been unable to reveal the homologies. These are proposed to be  
"hidden" in (alpha/beta)8-barrel enzymes. The term "hidden homology"  
means that the alignment of sequence stretches proposed to be homologous  
need not be structurally fully satisfactory. This is due to the very long  
evolutionary history of all (alpha/beta)8-barrels. This work identifies  
so-called hidden homology around the strand beta 2 that is flanked by  
loops containing invariant glycines and prolines in 17 different  
(alpha/beta)8-barrel enzymes, i.e., roughly in half of all currently known  
(alpha/beta)8-barrel proteins. The search was based on the idea that a  
conserved sequence region of an (alpha/beta)8-barrel enzyme should be more  
or less conserved also in the equivalent part of the structure of the  
other enzymes with this folding motif, given their mutual evolutionary  
relatedness. For this purpose, the sequence region around the  
well-conserved second beta-strand of alpha-amylase flanked by the  
invariant glycine and proline (56\_GFTAIWTP, **Aspergillus**  
**oryzae** alpha-amylase numbering), was used as the  
sequence-structural template. The proposal that the second beta-strand of

(alpha/beta)8-barrel fold is important from the evolutionary point of view is strongly supported by the increasing trend of the observed beta 2-strand structural similarity for the pairs of (alpha/beta)8-barrel enzymes: alpha-amylase and the alpha-subunit of **tryptophan synthase**, alpha-amylase and mandelate racemase, and alpha-amylase and cyclodextrin glycosyltransferase. This trend is also in agreement with the existing evolutionary division of the entire family of (alpha/beta)8-barrel proteins.

L11 ANSWER 2 OF 2 HCAPLUS COPYRIGHT 2004 ACS on STN  
 ACCESSION NUMBER: 2000:457232 HCAPLUS  
 DOCUMENT NUMBER: 133:85094  
 TITLE: Protein production in *Aspergillus* mutant cells that produce decreased levels of toxin  
 INVENTOR(S): Christensen, Bjorn Eggert; Mollgaard, Henrik; Kaasgaard, Svend; Lehmbeck, Jan  
 PATENT ASSIGNEE(S): Novo Nordisk A/S, Den.  
 SOURCE: PCT Int. Appl., 66 pp.  
 CODEN: PIXXD2  
 DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 1  
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2000039322	A1	20000706	WO 1999-DK726	19991222
W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				
RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG				
EP 1141371	A1	20011010	EP 1999-960956	19991222
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO				
JP 2002533133	T2	20021008	JP 2000-591212	19991222
US 6383781	B1	20020507	US 1999-472364	19991223
US 2002197682	A1	20021226	US 2002-99704	20020315
PRIORITY APPLN. INFO.:				
		DK 1998-1726	A	19981223
		DK 1999-745	A	19990527
		US 1999-117396P	P	19990127
		US 1999-139593P	P	19990617
		WO 1999-DK726	W	19991222
		US 1999-472364	A3	19991223

AB A method is provided for producing a polypeptide of interest by (a) cultivating a mutant of a parent *Aspergillus* cell, wherein (i) the mutant comprises a first nucleic acid sequence encoding the polypeptide and a second nucleic acid sequence comprising a modification of at least one of the genes responsible for the biosynthesis or secretion of at least one toxin, and (ii) the mutant produces less of the toxin than the parent *Aspergillus* cell when cultured under the same conditions; and (b) isolating the polypeptide from the culture medium. Also, mutants of *Aspergillus* cells are provided, as well as methods for obtaining the mutant cells. Thus, the dimethylallylcycloacetoacetyl-L-**tryptophan synthase** gene (DCAT-S) was characterized in *Aspergillus oryzae*. Since this enzyme is involved in biosynthesis of cyclopiazonic acid (CPA), CPA-neg. strains were constructed by modification of the DCAT-S gene, and improved of lipase and xylanase genes was obsd. in the CPA-neg. strains. Strains lacking genes *aflR* and *omtA*, involved in aflatoxin biosynthesis, also demonstrated improved heterologous protein prodn.

REFERENCE COUNT: 4 THERE ARE 4 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

=> d his

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FILE 'MEDLINE, HCAPLUS, BIOSIS, BIOTECHDS, SCISEARCH, EMBASE' ENTERED AT 16:56:45 ON 29 JUL 2004

L1 1 S DIMETHYLALLYL-CYCLOACETOACETYL-L-TRYPTOPHAN SYNTHASE

FILE 'REGISTRY' ENTERED AT 16:57:37 ON 29 JUL 2004

L2 0 S DIMETHYLALLYL-CYCLOACETOACETYL-L-TRYPTOPHAN SYNTHASE

L3 1 S DIMETHYLALLYL-CYCLOACETOACETYL TRYPTOPHAN SYNTHASE

L4 1 S DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE

FILE 'MEDLINE, HCAPLUS, BIOSIS, BIOTECHDS, SCISEARCH, EMBASE' ENTERED AT 16:59:46 ON 29 JUL 2004

L5 1 S DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE

L6 0 S DIMETHYLALLYLCYCLOACETOACETY SYNTHASE

L7 0 S DIMETHYLALLYL CYCLOACETOACETYL L-TRYPTOPHAN SYNTHASE

L8 125 S ASPERGILLUS ORYZAE AND SYNTHASE

L9 62 DUP REM L8 (63 DUPLICATES REMOVED)

L10 1 S L9 AND DIMETHYLALLYLCYCLOACETOACETYLTRYPTOPHAN SYNTHASE

L11 2 S L9 AND TRYPTOPHAN SYNTHASE

=> log y

COST IN U.S. DOLLARS	SINCE FILE ENTRY	TOTAL SESSION
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FULL ESTIMATED COST 16.45 79.30

DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)	SINCE FILE ENTRY	TOTAL SESSION
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L1: Entry 1 of 1

File: USPT

May 7, 2002

US-PAT-NO: 6383781DOCUMENT-IDENTIFIER: US 6383781 B1

TITLE: Methods for producing polypeptides in aspergillus mutant cells

DATE-ISSUED: May 7, 2002

**INVENTOR-INFORMATION:**

NAME	CITY	STATE	ZIP CODE	COUNTRY
Christensen; Bjorn Eggert	Bagsvaerd			DK
Mollgaard; Henrik	Lyngby			DK
Kaasgaard; Svend	Soborg			DK
Lehmbeck; Jan	Vekso			DK

**ASSIGNEE-INFORMATION:**

NAME	CITY	STATE	ZIP CODE	COUNTRY	TYPE CODE
Novozymes A/S	Bagsvaerd			DK	03

APPL-NO: 09/ 472364 [PALM]

DATE FILED: December 23, 1999

**PARENT-CASE:**

CROSS-REFERENCE TO RELATED APPLICATIONS This application claims benefit to U.S. provisional application No. 06/117,396 filed on Jan. 27, 1999, and U.S. provisional application No. 60/139,593 filed on Jun. 17, 1999, and claims foreign priority under 35 U.S.C. 119 to Danish application no. PA 1998 01726 filed on Dec. 23, 1998, Danish application no. DA 1999 00745 filed on May 27, 1999, the contents of which are fully incorporated herein by reference.

**FOREIGN-APPL-PRIORITY-DATA:**

COUNTRY	APPL-NO	APPL-DATE
DK	1998 01726	December 23, 1998
DK	1999 00745	May 27, 1999

INT-CL: [07] C12 P 21/06, C12 N 1/14

US-CL-ISSUED: 435/69.1, 435/71.1, 435/71.2, 435/256.1

US-CL-CURRENT: 435/69.1, 435/256.1, 435/71.1, 435/71.2

FIELD-OF-SEARCH: 435/69.1, 435/71.1, 435/71.2, 435/172.3, 435/320.1, 435/252.3, 435/256.1

**PRIOR-ART-DISCLOSED:**

U.S. PATENT DOCUMENTS

PAT-NO	ISSUE-DATE	PATENTEE-NAME	US-CL
<input type="checkbox"/> <u>5958727</u>	September 1999	Brody et al.	435/69.1

## FOREIGN PATENT DOCUMENTS

FOREIGN-PAT-NO	PUBN-DATE	COUNTRY	US-CL
1271068	January 1994	SU	
WO 95/15390	June 1995	WO	
WO 95/15391	June 1995	WO	

## OTHER PUBLICATIONS

Abstract of article by Tudzynski et al., Mol Gen Genet, vol. 261, pp. 133-141 (1999).

Abstract of Russian Patent No. SU 1271068 A1.

ART-UNIT: 1653

PRIMARY-EXAMINER: Carlson; Karen Cochrane

ATTY-AGENT-FIRM: Lambiris; Elias Garbell; Jason

## ABSTRACT:

A method is provided for producing a polypeptide of interest by (a) cultivating a mutant of a parent Aspergillus cell, wherein (i) the mutant comprises a first nucleic acid sequence encoding the polypeptide and a second nucleic acid sequence comprising a modification of at least one of the genes responsible for the biosynthesis or secretion of at least one toxin, and (ii) the mutant produces less of the toxin than the parent Aspergillus cell when cultured under the same conditions; and (b) isolating the polypeptide from the culture medium. Also, mutants of Aspergillus cells are provided, as well as methods for obtaining the mutant cells.

17 Claims, 2 Drawing figures

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L1: Entry 1 of 1

File: USPT

May 7, 2002

US-PAT-NO: 6383781DOCUMENT-IDENTIFIER: US 6383781 B1

TITLE: Methods for producing polypeptides in aspergillus mutant cells

DATE-ISSUED: May 7, 2002

## INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Christensen; Bjorn Eggert	Bagsvaerd			DK
Mollgaard; Henrik	Lyngby			DK
Kaasgaard; Svend	Soborg			DK
Lehmbeck; Jan	Vekso			DK

US-CL-CURRENT: 435/69.1; 435/256.1, 435/71.1, 435/71.2

## CLAIMS:

What is claimed is:

1. A method for producing a polypeptide, said method comprising:

(a) cultivating a mutant of a parent Aspergillus cell, wherein (i) the mutant comprises a nucleic acid sequence encoding said polypeptide, and (ii) the mutant produces less of at least one toxin selected from the group consisting of emodin, kojic acid, malformin, 3-nitropropionic acid, ochratoxins, and secalonic acids than the parent Aspergillus cell when cultured under the same conditions; and

(b) isolating the polypeptide from the culture medium.

2. The method of claim 1, wherein the mutant produces at least 90% less of the toxin than the parent Aspergillus cell when cultured under the same conditions.

3. The method of claim 1, wherein the toxin is emodin.

4. The method of claim 1, wherein the toxin is kojic acid.

5. The method of claim 1, wherein the toxin is malformin.

6. The method of claim 1, wherein the toxin is 3-nitropropionic acid.

7. The method of claim 1, wherein the toxin is an ochratoxin.

8. The method of claim 1, wherein the toxin is a secalonic acid.
9. The method of claim 1, wherein the mutant produces less of at least two said toxins than the parent Aspergillus cell when cultured under the same conditions.
10. The method of claim 1, wherein the mutant additionally produces less of an aflatoxin.
11. The method of claim 1, wherein the mutant additionally produces less of a cyclopiazonic acid.
12. The method of claim 1, wherein the parent Aspergillus cells is a cell from a subgroup selected from the group consisting of Chaetosartorya, Emericella, Eurotium, Fenellia, Hemicarpenteles, Neosartorya, Petromyces, Satoia, and Sclerocleista.
13. The method of claim 1, wherein the polypeptide of interest is native to the Aspergillus cell.
14. The method of claim 13, wherein the amount of the polypeptide produced by the mutant is greater than the amount produced by the parent Aspergillus cell when cultured under the same conditions.
15. The method of claim 1, wherein the polypeptide is heterologous to the mutant.
16. The method of claim 1, wherein the polypeptide is selected from the group consisting of a hormone or a precursor thereof, an enzyme or an enzyme variant or a precursor thereof, an antibody or a functional fragment thereof, a receptor or a functional fragment thereof, and a reporter.
17. The method of claim 16, wherein the polypeptide is selected from the group consisting of aminopeptidase, alpha-galactosidase, alpha-glucosidase, amylase, beta-galactosidase, beta-glucosidase, carbohydراse, carboxypeptidase, catalase, cellulase, chitinase, cutinase, cyclodextrin glycosyltransferase, deoxyribonuclease, endo-peptidase, exo-peptidase, esterase, galactanase, glucoamylase, invertase, laccase, lipase, lyase, mannase, mannosidase, mutanase, oxidase, oxygenase, pectate lyase, pectinase, peroxidase, phytase, polyphenoloxidase, protease, ribonuclease, transglutaminase, and xylanase.

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<a href="#">Generate OACS</a>				

Search Results - Record(s) 1 through 3 of 3 returned.

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1. Document ID: US 20020197682 A1

Using default format because multiple data bases are involved.

L1: Entry 1 of 3

File: PGPB

Dec 26, 2002

PGPUB-DOCUMENT-NUMBER: 20020197682  
 PGPUB-FILING-TYPE: new  
 DOCUMENT-IDENTIFIER: US 20020197682 A1

TITLE: Methods for producing polypeptides in Aspergillus mutant cells

PUBLICATION-DATE: December 26, 2002

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY	RULE-47
Christensen, Bjorn Eggert	Bagsvaerd		DK	
Mollgaard, Henrik	Lyngby		DK	
Kaasgaard, Svend	Soborg		DK	
Lehmbeck, Jan	Vekso		DK	

US-CL-CURRENT: 435/71.1; 435/254.3

<a href="#">Full</a>	<a href="#">Title</a>	<a href="#">Citation</a>	<a href="#">Front</a>	<a href="#">Review</a>	<a href="#">Classification</a>	<a href="#">Date</a>	<a href="#">Reference</a>	<a href="#">Sequences</a>	<a href="#">Attachments</a>	<a href="#">Claims</a>	<a href="#">KINIC</a>	<a href="#">Drawn D</a>
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2. Document ID: US 6383781 B1

L1: Entry 2 of 3

File: USPT

May 7, 2002

US-PAT-NO: 6383781  
 DOCUMENT-IDENTIFIER: US 6383781 B1

TITLE: Methods for producing polypeptides in aspergillus mutant cells

DATE-ISSUED: May 7, 2002

INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Christensen; Bjorn Eggert	Bagsvaerd			DK
Mollgaard; Henrik	Lyngby			DK
Kaasgaard; Svend	Soborg			DK
Lehmbeck; Jan	Vekso			DK

US-CL-CURRENT: 435/69.1; 435/256.1, 435/71.1, 435/71.2

Full	Title	Citation	Front	Review	Classification	Date	Reference	Claims	KINIC	Drawn D
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3. Document ID: US 20020197682 A1, WO 200039322 A1, AU 200017740 A, EP 1141371 A1, CN 1333837 A, US 6383781 B1, JP 2002533133 W

L1: Entry 3 of 3

File: DWPI

Dec 26, 2002

DERWENT-ACC-NO: 2000-452411

DERWENT-WEEK: 200304

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TITLE: Producing a polypeptide of interest such as a hormone or enzyme, comprising cultivating a mutant of a parent Aspergillus cell which produces less of at least one toxin of interest compared to the parent cell under the same conditions

INVENTOR: CHRISTENSEN, B E; KAASGAARD, S ; LEHMBECK, J ; MOLLGAARD, H

PRIORITY-DATA: 1999DK-0000745 (May 27, 1999), 1998DK-0001726 (December 23, 1998)

## PATENT-FAMILY:

PUB-NO	PUB-DATE	LANGUAGE	PAGES	MAIN-IPC
<u>US 20020197682 A1</u>	December 26, 2002		000	C12P021/02
<u>WO 200039322 A1</u>	July 6, 2000	E	061	C12P021/00
<u>AU 200017740 A</u>	July 31, 2000		000	C12P021/00
<u>EP 1141371 A1</u>	October 10, 2001	E	000	C12P021/00
<u>CN 1333837 A</u>	January 30, 2002		000	C12P021/00
<u>US 6383781 B1</u>	May 7, 2002		000	C12P021/06
<u>JP 2002533133 W</u>	October 8, 2002		061	C12N015/09

INT-CL (IPC): C12 N 1/14; C12 N 1/15; C12 N 1/16; C12 N 9/00; C12 N 9/10; C12 N 9/20; C12 N 9/42; C12 N 9/10; C12 N 15/01; C12 N 15/09; C12 P 21/00; C12 P 21/02; C12 P 21/06; C12 P 21/00; C12 R 1:66; C12 N 1/15; C12 R 1:69; C12 R 1:66; C12 R 1:66; C12 P 21/00; C12 N 9/10

Full	Title	Citation	Front	Review	Classification	Date	Reference	Claims	KINIC	Drawn D
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Terms	Documents
dimethylallyl-cycloacetoacetyl-L-tryptophan synthase	3

Display Format: [-] 

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## WEST Search History

DATE: Thursday, July 29, 2004

<u>Hide?</u>	<u>Set Name</u>	<u>Query</u>	<u>Hit Count</u>
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<input type="checkbox"/>	L4	L2 and synthase	580
<input type="checkbox"/>	L3	L2 with synthase	0
<input type="checkbox"/>	L2	435/193.ccls.	1906
<input type="checkbox"/>	L1	dimethylallyl-cycloacetoacetyl-L-tryptophan synthase	3

END OF SEARCH HISTORY